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Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866)  
217-9197 (toll free).

Reviewer: Keisha Douglas

Timestamp: [year=2008; month=8; day=5; hr=16; min=35; sec=37; ms=897; ]

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Reviewer Comments:

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<211> 1588

<212> DNA

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<221> CDS

<222> (1)...(1059)

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Per the above sequence id# 2 is invalid, please insert (3) letter  
codings for amino acids immediately below  
corresponding nucleotide triplets. This error is seen globally  
throughout the sequence.

<210> 5

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<213> Artificial Sequence

<220>

<223> Ligand peptide

<210> 9

<211> 4

<212> PRT

<213> Artificial Sequence

<220>

<223> added peptide

Per the above, both <223> responses for sequence id#'s 5 and 9 are invalid, please explain Artificial.

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Application No: 10785230 Version No: 2.0

Input Set:

Output Set:

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Finished: 2008-06-26 16:28:55.423  
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 893 ms  
Total Warnings: 12  
Total Errors: 0  
No. of SeqIDs Defined: 12  
Actual SeqID Count: 12

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# SEQUENCE LISTING

<110> KISHIMOTO, Tadimitsu  
NAGASAWA, Takashi  
TACHIBANA, Kazunobu

<120> Inhibiting vascularization using antibodies to CXCR4 and SDF-1

<130> 46124-5042-US01

<140> 10785230

<141> 2004-02-25

<150> US 09/646,785

<151> 2001-02-16

<150> PCT/JP99/01448

<151> 1999-03-23

<150> JP10/95448

<151> 1998-03-24

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 <222> (1)...(1059)

<400> 2

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tcc atc atc ttc tta act ggc att gtg ggc aat gga ttg gtc atc 180
ctg gtc atg ggt tac cag aag aaa ctg aga agc atg acg gac aag 225
tac agg ctg cac ctg tca gtg gcc gac ctc ctc ttt gtc atc acg 270
ctt ccc ttc tgg gca gtt gat gcc gtg gca aac tgg tac ttt ggg 315
aac ttc cta tgc aag gca gtc cat gtc atc tac aca gtc aac ctc 360
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Ser	Ser	Leu	Lys	Ile	Leu	Ser	Lys	Gly	Lys	Arg	Gly	Gly	His	Ser	
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gac aag tac cgg ctg cac ctg tca gtg gct gac ctc ctc ttt gtc 270
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cag ggg gac atc agt cag ggg gat gac agg tac atc tgt gac cgc 585
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ata atg gtg ggt ctc atc ctg ccc ggc atc gtc atc ctc tcc tgt 675
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<213> Artificial Sequence

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Pro	Cys	Arg	Phe	Phe	Glu	Ser	His	Val	Ala	Arg	Ala	Asn	Val	Lys
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cca tgc cga ttc ttc gaa agc cat gtt gcc aga gcc aac gtc aag						608
cat ctc aaa att ctc aac act cca aac tgt gcc ctt cag att gta						653
gcc cgg ctg aag aac aac aac aga caa gtg tgc att gac ccg aag						698
cta aag tgg att cag gag tac ctg gag aaa gct tta aac aag taa						743
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<212> PRT
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          35              40              45
His Leu Lys Ile Leu Asn Thr Pro Asn Cys Ala Leu Gln Ile Val
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Ala Arg Leu Lys Asn Asn Arg Gln Val Cys Ile Asp Pro Lys
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<212> DNA
<213> Mus

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<222> (82)...(351)

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81

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cat ctg aaa atc ctc aac act cca aac tgt gcc ctt cag att gtt 261  
gca cgg ctg aag aac aac aac aga caa gtg tgc att gac ccg aaa 306  
tta aag tgg atc caa gag tac ctg gag aaa gct tta aac aag taa 351  
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